

IFWO

RAW SEQUENCE LISTING

1 <110> APPLICANT: GESTRELIUS, STINA 2 HAMMARSTROM, LARS DATE: 09/17/2004 TIME: 09:06:46

PATENT APPLICATION: US/10/698,618

Input Set : N:\Crf3\RULE60\10698618.raw.txt
Output Set: N:\CRF4\09172004\J698618.raw

```
LYNGSTADAAS, PETTER
        ANDERSSON, CHRISTER
 5
        SLABY, IVAN
        HAMMARGREN, TOMAS
 7 <120> TITLE OF INVENTION: MATRIX PROTEIN COMPOSITIONS FOR WOUND HEALING
 8 <130> FILE REFERENCE: 47927-48292-CPA
 9 <140> CURRENT APPLICATION NUMBER: US/10/698,618
10 <141> CURRENT FILING DATE: 2003-10-30
11 <150> PRIOR APPLICATION NUMBER: US/10/156,300
12 <151> PRIOR FILING DATE: 2002-05-28
13 <150> PRIOR APPLICATION NUMBER: US/09/258,613
                                                      ENTERED
14 <151> PRIOR FILING DATE: 1999-02-26
15 <150> PRIOR APPLICATION NUMBER: DK PA 1998 01328
16 <151> PRIOR FILING DATE: 1998-10-16
                                                 17 <150> PRIOR APPLICATION NUMBER: 60/081,551
18 <151> PRIOR FILING DATE: 1998-04-13
19 <150> PRIOR APPLICATION NUMBER: DK 0270/98
20 <151> PRIOR FILING DATE: 1998-02-27
21 <160> NUMBER OF SEQ ID NOS: 2
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 407
26 <212> TYPE: PRT
27 <213> ORGANISM: Rattus sp.
28 <400> SEQUENCE: 1
29
        Met Ser Ala Ser Lys Ile Pro Leu Phe Lys Met Lys Gly Leu Leu Leu
30
31
        Phe Leu Ser Leu Val Lys Met Ser Leu Ala Val Pro Ala Phe Pro Gln
32
                                         25
33
        Arg Pro Gly Gly Gln Gly Met Ala Pro Pro Gly Met Ala Ser Leu Ser
34
35
        Leu Glu Thr Met Arg Gln Leu Gly Ser Leu Gln Gly Leu Asn Ala Leu
36
                                 55
37
        Ser Gln Tyr Ser Arg Leu Gly Phe Gly Lys Ala Leu Asn Ser Leu Trp
38
                                                75
39
        Leu His Gly Leu Leu Pro Pro His Asn Ser Phe Pro Trp Ile Gly Pro
40
                         85
                                            90
41
        Arg Glu His Glu Thr Gln Gln Pro Ser Leu Gln Pro His Gln Pro Gly
42
                                       105
43
        Leu Lys Pro Phe Leu Gln Pro Thr Ala Ala Thr Gly Val Gln Val Thr
44
                115
                                   120
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45
          Pro Gln Lys Pro Gly Pro His Pro Pro Met His Pro Gly Gln Leu Pro
46
              130
                                                       140
          Leu Gln Glu Gly Glu Leu Ile Ala Pro Asp Glu Pro Gln Val Ala Pro
47
48
                              150
49
          Ser Glu Asn Pro Pro Thr Pro Glu Val Pro Ile Met Asp Phe Gly Asp
50
                          165
                                               170
          Pro Gln Phe Pro Thr Val Phe Gln Ile Ala His Ser Leu Ser Arg Gly
51
52
                      180
                                           185
53
         Pro Met Ala His Asn Lys Val Pro Thr Phe Tyr Pro Gly Met Phe Tyr
54
                                      200
         Met Ser Tyr Gly Ala Asn Gln Leu Asn Ala Pro Gly Arg Ile Gly Phe
55
56
                                                       220
57
         Met Ser Ser Glu Glu Met Pro Gly Glu Arg Gly Ser Pro Met Gly Tyr
58
                              230
                                                   235
59
         Gly Thr Leu Phe Pro Gly Tyr Gly Gly Phe Arg Gln Thr Leu Arg Gly
60
                                               250
61
         Leu Asn Gln Asn Ser Pro Lys Gly Gly Asp Phe Thr Val Glu Val Asp
62
                                           265
63
         Ser Pro Val Ser Val Thr Lys Gly Pro Glu Lys Gly Glu Gly Pro Glu
64
                                      280
                                                           285
         Gly Ser Pro Leu Gln Glu Pro Ser Pro Asp Lys Gly Glu Asn Pro Ala
65
66
                                  295
67
         Leu Leu Ser Gln Ile Ala Pro Gly Ala His Ala Gly Leu Leu Ala Phe
68
                                                   315
         Pro Asn Asp His Ile Pro Asn Met Ala Arg Gly Pro Ala Gly Gln Arg
69
70
                          325
71
         Leu Leu Gly Val Thr Pro Ala Ala Ala Asp Pro Leu Ile Thr Pro Glu
72
                                          345
         Leu Ala Glu Val Tyr Glu Thr Tyr Gly Ala Asp Val Thr Thr Pro Leu
73
74
                                      360
75
         Gly Asp Gly Glu Ala Thr Met Asp Ile Thr Met Ser Pro Asp Thr Gln
76
                                  375
77
         Gln Pro Pro Met Pro Gly Asn Lys Val His Gln Pro Gln Val His Asn
78
                              390
                                                  395
79
         Ala Trp Arg Phe Gln Glu Pro
80
                         405
82 <210> SEQ ID NO: 2
83 <211> LENGTH: 324
84 <212> TYPE: PRT
85 <213> ORGANISM: Rattus sp.
86 <400> SEQUENCE: 2
         Met Lys Pro Asn Ser Met Glu Asn Ser Leu Pro Val His Pro Pro
87
88
                           5
                                               10
         Leu Pro Ser Gln Pro Ser Leu Gln Pro His Gln Pro Gly Leu Lys Pro
89
90
                                           25
91
         Phe Leu Gln Pro Thr Ala Ala Thr Gly Val Gln Val Thr Pro Gln Lys
92
                                       40
93
         Pro Gly Pro His Pro Pro Met His Pro Gly Gln Leu Pro Leu Gln Glu
94
                                  55
```

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95 96	Gly 65	Glu	Leu	Ile	Ala	Pro 70	Asp	Glu	Pro	Gln	Val 75	Ala	Pro	Ser	Glu	
97		Pro	Thr	Pro	G111		Dro	т1д	Mot	λan	-	C1	7 00	Deca	a1	80
98					85	vui	110	116	Mec	90 90	FIIG	GIY	Asp	PIO	95	Pne
99	Pro	Thr	Val	Phe		Tle	Δla	His	Ser		Ser	Δνα	Gl _V	Dro		7.7.
100				100					105		DCI	Arg	Сту	110		Ата
101	His	Asn	Lvs	-		Thr	Phe	Tvr			, Met	Phe	Tur			Tyr
102			115					120		. 017			125		. DCI	TYT
103	Gly	Ala	Asn	Gln	Leu	Asn	Ala			Arc	r Ile	G]v			Ser	Ser
104	_	130					135		2		,	140			001	OCI
105	Glu	Glu	Met	Pro	Gly	Glu	Arq	Gly	Ser	Pro	Met			Glv	Thr	Leu
106	145				-	150		_			155		- 1 -	0-1		160
107	Phe	Pro	Gly	Tyr	Gly	Gly	Phe	Arg	Gln	Thr	Leu	Arq	Gly	Leu	Asn	Gln
108					165			_		170			•		175	
109	Asn	Ser	Pro	Lys	Gly	Gly	Asp	Phe	Thr	Val	Glu	Val	Asp	Ser	Pro	Val
110				180					185					190		
111	Ser	Val	Thr	Lys	Gly	Pro	Glu	Lys	Gly	Glu	Gly	Pro	Glu	Gly	Ser	Pro
112			195					200					205			
113	Leu	Gln	Glu	Pro	Ser	Pro	Asp	Lys	Gly	Glu	Asn	Pro	Ala	Leu	Leu	Ser
114		210					215					220				
115	Gln	Ile	Ala	Pro	Gly		His	Ala	Gly	Leu	Leu	Ala	Phe	Pro	Asn	Asp
116	225					230					235					240
117	His	Ile	Pro	Asn		Ala	Arg	Gly	Pro	Ala	Gly	Gln	Arg	Leu	Leu	Gly
118				_	245					250					255	
119	Val	Thr	Pro		Ala	Ala	Asp	Pro		Ile	Thr	Pro	Glu	Leu	Ala	Glu
120			_	260					265					270		
121	Val	Tyr	Glu	Thr	\mathtt{Tyr}	Gly	Ala		Val	Thr	Thr	Pro	Leu	Gly	Asp	Gly
122			275			_		280					285			
123	GIu	Ala	Thr	Met	Asp	Ile		Met	Ser	Pro	Asp	Thr	Gln	Gln	Pro	Pro
124		290					295					300				
125	Met	Pro	Gly	Asn	Lys		His	Gln	Pro	Gln		His	Asn	Ala	\mathtt{Trp}	Arg
126	305	~1	~1	_		310					315					320
127	Pne	GIn	Glu	Pro												

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\10698618.raw.txt Output Set: N:\CRF4\09172004\J698618.raw